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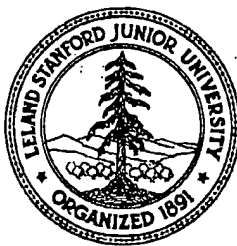
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Exhibit A

# LABORATORY NOTEBOOK



Department Chemical Engineering  
Stanford University  
Stanford, California

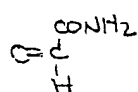
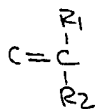
*Ying Chang*

*(650) 7239140*

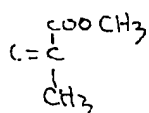
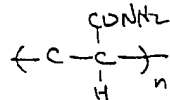
*Rm 105 Stauffer II*

acrylic family (Principles of polymerization 4, Odian 3rd ed. Wiley Interscience)

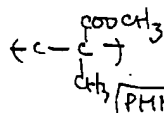
(P. 311)



acrylamide



methyl methacrylate  
(MMA)



PMMA

Process:  
solution, suspension  
emulsion

R<sub>1</sub> COOH

COOH

CN

Cl

R<sub>2</sub> H

CH<sub>3</sub>

H

Cl

name poly(acrylic acid)  
water soluble

poly(methacrylic acid)  
"

polyacrylonitrile

poly(vinylidene chloride)

R<sub>1</sub> -OCOCH<sub>3</sub>

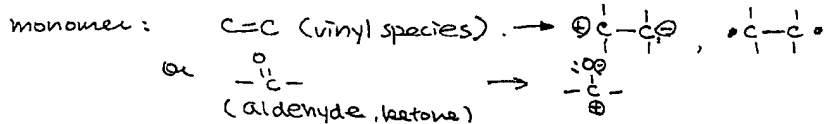
R<sub>2</sub> -H

name poly(vinyl acetate)

water based paint

radical  
ionic polymerization

chain polymerization



type of chain polymerization undergone by various unsat. monomers

monomer

type of initiation

radical

cationic

anionic

ethylene

+

-

+

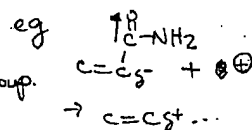
vinyl ester, acrylate,  
methacrylate, acrylonitrile,  
methacrylonitrile

+

+

-

electron  
withdrawn group



styrene, methylstyrene

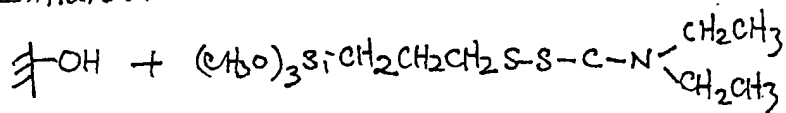
+

+

+

Ref: L. Liang, X. Feng, J. Liu P.C. Rieke, G.E. Fryxell Macromolecules 1998 31 7845-50.

Initiator:

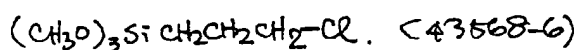
N,N'-diethylamino dithiocarbamoyl  
propyl(trimethoxy)silane

DAPMS

Ref 15, 16.

+  
NIPAAm/BisAAM  
UV, 254 nm

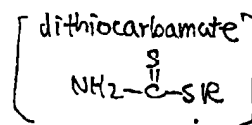
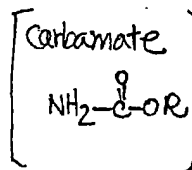
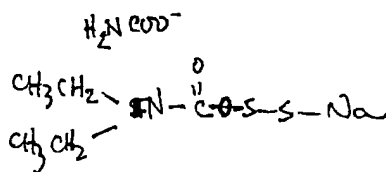
Reaction

Reflux  
48h 56°C250ml 3-neck flask  
12g CPTMS + 4g SPDC + 100ml acetone  
3-chloropropyltrimethoxysilane, 97% Aldrich → purified reduced distillation.

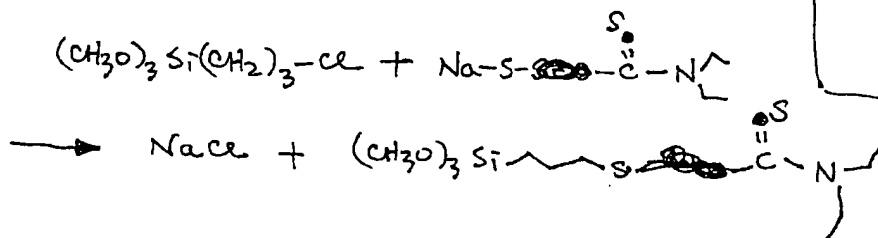
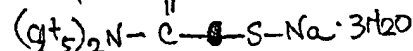
3-chloropropyl triethoxysilane 95% Aldrich (108-2)  
[5089 7033 SIC 24070]

SPDC:

Sodium, N'-diethyl dithiocarbamate



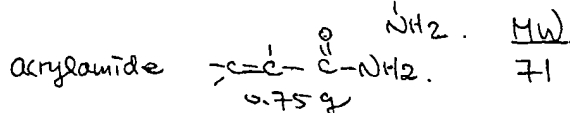
Reaction:

CP: sodium diethyl dithiocarbamate  
trihydrate

FW = 225.31

mp = 95-98.5°

hygroscopic. 22868-0 5g 17.55

$$\frac{1}{0.075} = \frac{1000}{75} = 13.$$


$$\frac{0.75g}{71} \sim 0.01 \text{ mol.}$$

$$0.01 \text{ mol acrylamide} / 0.75 \text{ ml.}$$

2.2' dipyrdayl  $0.036$   $156.19$   $(2.3 \times 10^{-4} \text{ mol}) = 0.01 \text{ mol}$   $0.01 \text{ mol}$   $0.75 \times 10^{-3} \text{ L} = 1.3 \text{ M}$   $13 \text{ M}$   $\text{too high!}$

0.0067

One  
0.75

$$\text{Cu(I)Cl} : \text{bipy} = 1 : 3 \quad (\text{~~10}^{-4} \text{ mol})~~$$
  

$$= 10^{-4} : 3 \times 10^{-4} \text{ (mol)}$$

Calc. ( )

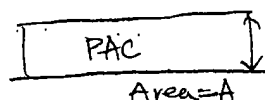
(Wang, 1995)

$$M_0: Cu(I)Cl$$

$$= 100:1$$

$$\begin{aligned} &= 0.01 g : 3 \times 156.19 \times 10^{-4} = 10^{-2} \text{ mol} : 10^{-4} \text{ mol} \\ &= 0.01 g : 0.04685 g = 0.71 : 0.01 g \end{aligned}$$

What thickness means in this case



Assumption: density is const = bulk material

$d_4^{20} = 1.189$  For PAC 50% wt in  $H_2O$ .

$$1.189 = \frac{1+x}{2} \Rightarrow d_{PAC} \sim 1.37$$

$$\left( \text{needed mole of monomers} \right) = \frac{\Delta h d (\text{total weight})}{71} = \frac{1 \text{ cm}^2 \times 1.37}{71} \text{ h(cm)} = 0.019 \times 10^{-7} \text{ h(cm)} \\ \cong 2 \times 10^{-9} \text{ h(cm)} \text{ moles.}$$

According to this, we can greatly reduce the amt of monomers from original recipe

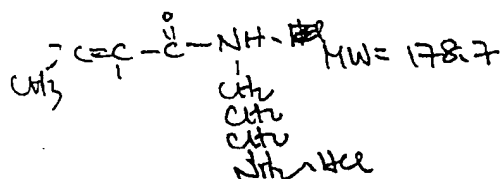
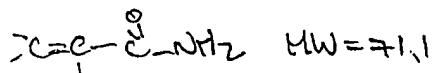
2x:  $h = 10^4 \text{ nm} (= 10 \mu\text{m})$   
 $\Rightarrow \text{needed monomers} = 2 \times 10^{-5} \text{ mol}$

Assume densely packed. 1 molecule/nm<sup>2</sup>.  
 For A = 1 cm<sup>2</sup> = 10<sup>14</sup> nm<sup>2</sup> ⇒ 10<sup>14</sup> ~~molecules~~ # PAC =  $\frac{10^{14}}{6 \times 10^{23}} = 1.7 \times 10^{-10}$   
 DP =  $2 \times 10^{-5}$  mol /  $2 \times 10^{-10}$  = 10<sup>5</sup> 10<sup>5</sup> × 0.5 = 5 × 10<sup>4</sup> nm ~ 0.1 (10 μm)  
 (good agreement!)

For DMF ~~10~~ 1 ml. make 0.01 M acrylamide soln =  $0.01 \text{ mol} \times 71 = 0.71 \text{ g}$   
 $0.01 = \frac{x \text{ mol}}{10^{-3} \text{ l}} \Rightarrow x = 10^{-5} \text{ mol} \cdot 10^{-5} \text{ mol} \times 71 = 0.00071 \text{ g}$

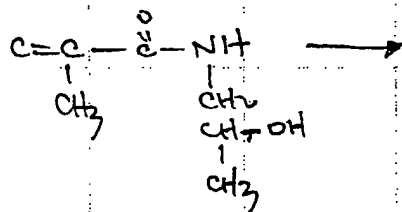
Roughly for DMF 10 ml acrylamide  $0.0071 \text{ g} \Rightarrow \frac{0.0001}{0.01} = 10^{-2} \text{ M}$ .

Better soln:  $0.1\text{ M} \Rightarrow 10\text{ mL DMF} + 0.071\text{ g acrylamide}$  good enough for cover the whole surface.



Signed..... Date.....

Objective: Repeat protocol exp 7. For N-(2-hydroxypropyl) ~~acryl~~ methacrylamide



HIV6d N-(2-hydroxypropyl) methacrylamide [21442-01-3] MW 143.2 mp = 67° 10g 12/25

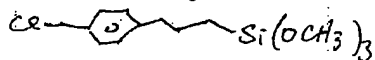
$\text{H}_2\text{C}=\text{C}-\text{CH}_3$  FW = 130.14 10M  
 $\text{COOCH}_2\text{CH}_2\text{OH}$  d = 1.073 HEMA in 10ml DMF 0.1 mol = 13g HEMA

+  
 $\text{Cu(I)Cl}(\text{bipyridyl})_2$  CuCl<sub>2</sub> (MW = 99) 0.23g (0.0023 mol) 1  
 +  
 bipyridyl ~~0.23g~~ (MW = 156.19) 1.08g (0.0069 mol) 3



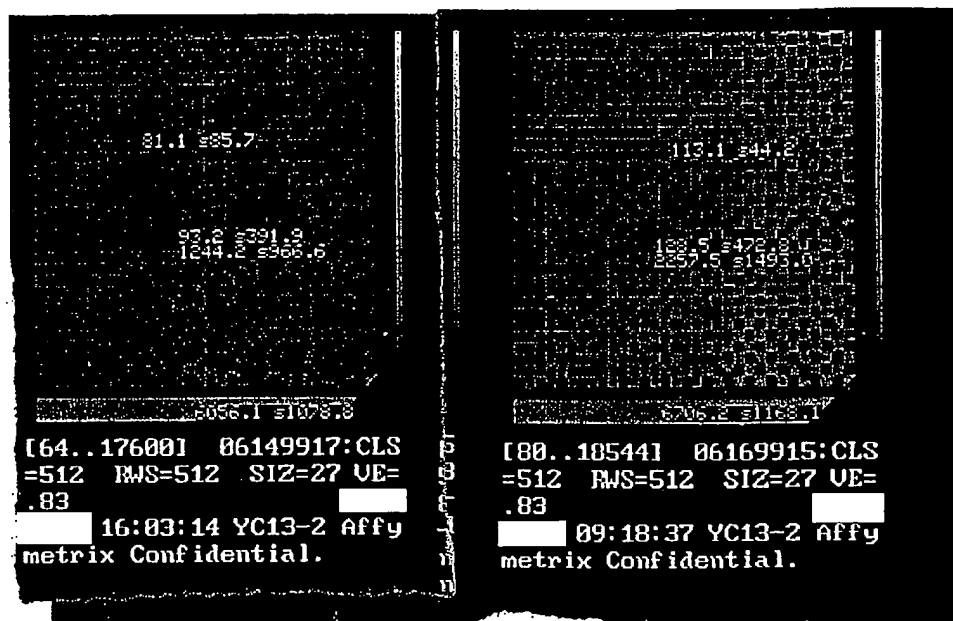
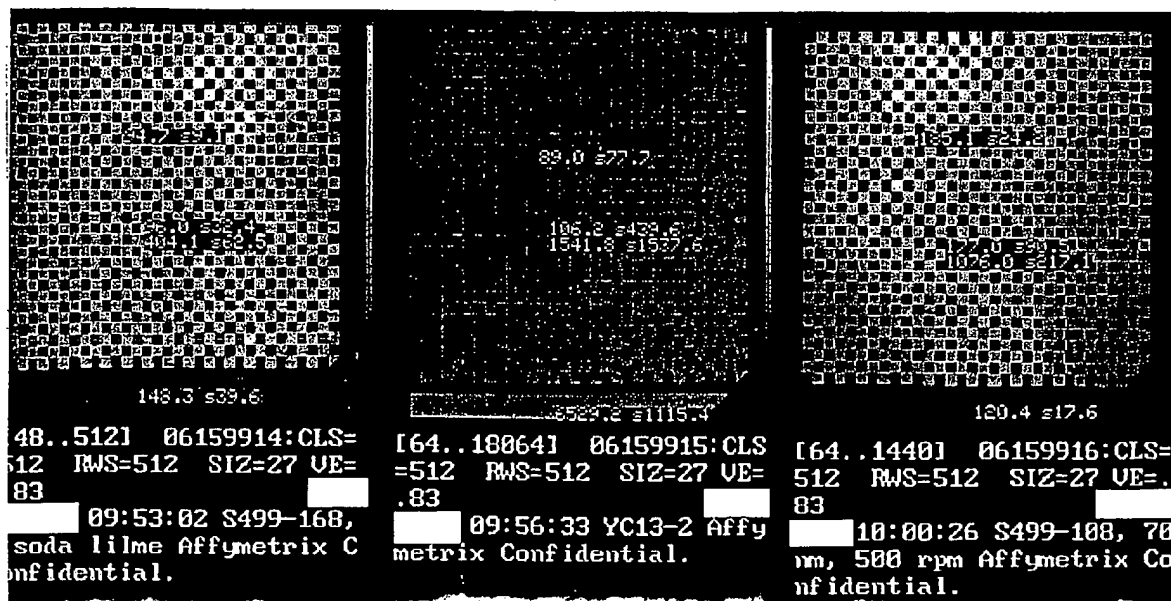
new materials.

chlorophenyl propyl trimethoxysilane 30min 1% in acetone.



+

↓ 1h

↑  
17h↑  
40h

Objective: AIBN — HEMA.

glass — 10 pieces. Si substrates — 4 pieces.

Comment: last exp the thickness only reaches up to 10 nm. I think it's due to the low surface coverage of AIBN layer (I forgot to work it in darkness, so the final thickness checked by Si wafer sample is only half of the thickness of exp 10.)

Improvement: This time I will work under dark and high concn of initiator layer.

Sample	Initiator Condition	HEMA rxn.	deprotection	hybridization
13-1	glass + AIBN/ toluene dark, E <sub>2</sub> N, rt temp in N <sub>2</sub> glove bag.	DNA strands on pattern work on condition 210-3 20 to HEMA/DMF 65°C 24h. degass. freeze thaw 3 times HEMA 40ml DMF 160ml store in darkness.	EtOH/EDA ethylene diamine (1/1) 4h → w/ DI H <sub>2</sub> O	500 M. DNA target - tag in 6XSSPE soln (1h) shake rt temp. ↓ 17 hr. (hyb) scan 10am 1st scan @ ↓ 24 hr (total 40h) (hyb) 2nd scan ↓ not scan
13-2	starting			another protocol. 45°C 16h.
13-3	4:00 pm			
13-4	~			
13-5				
13-6				
13-7				
13-8				

Signed \_\_\_\_\_ Date \_\_\_\_\_

hyl

sample	hyp time(h)	hyp	backgd	fluorescein	hyp/bg	fluor/bg
e13-2	1	1244	88	6056	14.1	68.8
control 1	1					
control 2	1	362	66	171	5.5	2.6
e13-2/control		3	1	35	3	27

1244/362 88/66 6056/171 14.1/5.5 68.8/2.6

sample	hyp time(h)	hyp	backgd	fluorescein	hyp/bg	fluore/bg
e13-2	17	1542	89	6529	17.3	73.4
control 1	17	339	101	140	3.4	1.4
control 2	17	404	95	148	4.3	1.6
e13-2/control		4	1	45	5	50

sample	hyp time(h)	hyp	backgd	fluorescein	hyp/bg	fluore/bg
e13-2	40	2258	113	6706.2	20.0	59.3
control 1	40	225	94	99	2.4	1.1
control 2	40	318	93	114	3.4	1.2
e13-2/control		8	1	63	7	52

experiment 13

report on file: e13-fluor.xls (thin's chang)

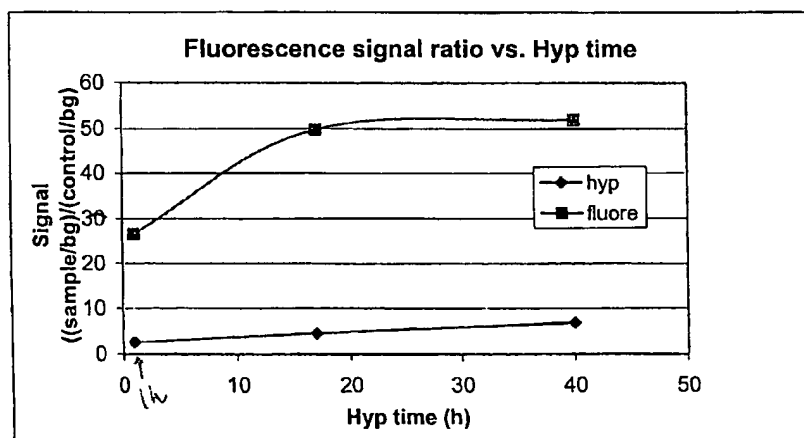
pHEMA initiated by AIBN

Fluorescence measured by Marc Glazer

Experiment condition: AIBN(20h)- pHEMA(20h 65c) - DNA probes-hybridization

estimated pHEMA thickness: 5 nm (estimated by Si(100) ellipsometry; need further confirmed by XPS)

Comments for improvement: high standard deviation(ca 1500) -- surface film is not uniform enough -- ne  
long hybridization time



Signed

S. C. Chang

Date

Subject.....

rxn  
condition

13-9

"

(same as  
~~last~~  
previous)

13-10

"

13-11

same  
batch  
but  
Si(100)

13-12

Si(100)

13-13

"

13-14

"

Signed..... Date.....